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PATENT APPLICATION: US/09/308,435A   TIME: 11:52:03

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Output Set: N:\CRF3\04262001\I308435A.raw

Does Not Comply
Corrected Diskette Needed
P.6

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 5 <120> TITLE OF INVENTION: Vaccine Delivery System and Method of Production
 7 <130> FILE REFERENCE: 1103326-0560
 9 <140> CURRENT APPLICATION NUMBER: 09/308,435A
10 <141> CURRENT FILING DATE: 1999-05-19
12 <150> PRIOR APPLICATION NUMBER: PCT/SE99/00582
13 <151> PRIOR FILING DATE: 1999-04-09
15 <150> PRIOR APPLICATION NUMBER: SE 9801288-3
16 <151> PRIOR FILING DATE: 1998-04-14
18 <160> NUMBER OF SEQ ID NOS: 25
20 <170> SOFTWARE: PatentIn Ver. 2.1
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24 <212> TYPE: DNA
25 <213> ORGANISM: Helicobacter pylori
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28 <221> NAME/KEY: CDS
29 <222> LOCATION: (793)..(1572)
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36 gatttgattt tagggaatta catgcaagtg aatgaaaaaa acattcaagc gtttgccccc 180
38 aaacaataag gtaaaaaatg ccactcactc atttgaatga agaaaatcaa cctaaaatgg 240
40 tggatatagg ggataaagaa accactgaaa gaatcgctct agcaagcggt cgtatcagca 300
42 tgaataaaga ggcttatgac gctattatca atcatggcgt caaaaagggt ccggtattac 360
44 aaactgctat tattgctggg attatggggg ctaaaaagac aagcgaactc attcccatgt 420
46 gccatccaat catgctcaat ggggtggata ttgatatttt agaagaaaaa gagacttgta 480
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50 taatgagtgt gagcgtaggg cttttaacca tttatgacat ggtgaaagcc attgataaga 600
52 gcatgacaat tagcggtgtg atgctggaat ataaaagtgg aggcaaaagt ggggattata 660
54 acgctaaaaa atagaaaaag actgataatc taaagatatt aggtaaaat aacattttga 720
56 caacaaaagc gtgttggttg cttcggattt gttgttatag aagtctaaaa tattacaatc 780
58 aaggatagaa cg atg aga gca aat aat cat ttt aaa gat ttt gca tgg aaa 831
59                Met Arg Ala Asn Asn His Phe Lys Asp Phe Ala Trp Lys
60                  1               5                  10
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63 Lys Cys Leu Leu Gly Ala Ser Val Val Ala Leu Leu Val Gly Cys Ser
64     15              20                  25
66 ccg cat att att gaa acc aat gaa gtc gct ttg aaa ttg aat tac cat  927
67 Pro His Ile Ile Glu Thr Asn Glu Val Ala Leu Lys Leu Asn Tyr His
68  30              35                  40                  45
70 cca gct agc gag aaa gtt caa gcg tta gat gaa aag att ttg ctt tta  975
71 Pro Ala Ser Glu Lys Val Gln Ala Leu Asp Glu Lys Ile Leu Leu Leu
72                  50                  55                  60
74 agg cca gct ttc caa tat agc gat aat atc gct aaa gag tat gaa aac 1023
75 Arg Pro Ala Phe Gln Tyr Ser Asp Asn Ile Ala Lys Glu Tyr Glu Asn
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83 Gln Gly Tyr Lys Val Ile Ser Val Asp Ser Ser Asp Lys Asp Asp Phe
84      95                  100                 105
86 tct ttt gca caa aaa aaa gaa ggg tat ttg gcg gtt gct atg aat ggc   1167
87 Ser Phe Ala Gln Lys Lys Glu Gly Tyr Leu Ala Val Ala Met Asn Gly
88 110                 115                 120                 125
90 gaa att gtt tta cgc ccc gat cct aaa agg acc ata cag aaa aaa tca   1215
91 Glu Ile Val Leu Arg Pro Asp Pro Lys Arg Thr Ile Gln Lys Lys Ser
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94 gaa ccc ggg tta tta ttc tcc acc ggt ttg gac aaa atg gaa ggg gtt   1263
95 Glu Pro Gly Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val
96          145              ·  150                 155
98 tta atc ccg gct ggg ttt att aag gtt acc ata cta gag cct atg agt   1311
99 Leu Ile Pro Ala Gly Phe Ile Lys Val Thr Ile Leu Glu Pro Met Ser
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102 ggg gaa tct ttg gat tct ttt acg atg gat ttg agc gag ttg gac att   1359
103 Gly Glu Ser Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp Ile
104      175                 180                 185
106 caa gaa aaa ttc tta aaa acc acc cat tca agc cat agc ggg ggg tta   1407
107 Gln Glu Lys Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly Leu
108 190                 195                 200                 205
110 gtt agc act atg gtt aag gga acg gat aat tct aat gac gcg atc aag   1455
111 Val Ser Thr Met Val Lys Gly Thr Asp Asn Ser Asn Asp Ala Ile Lys
112                  210                 215                 220
114 agc gct ttg aat aag att ttt gca aat atc atg caa gaa ata gac aaa   1503
115 Ser Ala Leu Asn Lys Ile Phe Ala Asn Ile Met Gln Glu Ile Asp Lys
116              225                 230                 235
118 aaa ctc act caa aag aat tta gaa tct tat caa aaa gac gcc aaa gaa   1551
119 Lys Leu Thr Gln Lys Asn Leu Glu Ser Tyr Gln Lys Asp Ala Lys Glu
120          240                 245                 ·250
122 tta aaa ggc aaa aga aac cga taaaaacaaa taacgcataa gaaaagaacg     1602
123 Leu Lys Gly Lys Arg Asn Arg
124      255                 260
126 cttgaataaa ctgcttaaaa agggtttttt agcgttcttt ttgagcgtgt atttaagggc 1662
128 tgatgatc                                                          1670
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133 <212> TYPE: PRT
134 <213> ORGANISM: Helicobacter pylori
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141              20                  25                  30
143 Ile Glu Thr Asn Glu Val Ala Leu Lys Leu Asn Tyr His Pro Ala Ser
144          35                  40                  45
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153                 85                  90                  95
155 Lys Val Ile Ser Val Asp Ser Ser Asp Lys Asp Asp Phe Ser Phe Ala
156                 100                 105                 110
158 Gln Lys Lys Glu Gly Tyr Leu Ala Val Ala Met Asn Gly Glu Ile Val
159         115                 120                 125
161 Leu Arg Pro Asp Pro Lys Arg Thr Ile Gln Lys Lys Ser Glu Pro Gly
162     130                 135                 140
164 Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val Leu Ile Pro
165 145                 150                 155                 160
167 Ala Gly Phe Ile Lys Val Thr Ile Leu Glu Pro Met Ser Gly Glu Ser
168                 165                 170                 175
170 Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp Ile Gln Glu Lys
171             180                 185                 190
173 Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly Leu Val Ser Thr
174         195                 200                 205
176 Met Val Lys Gly Thr Asp Asn Ser Asn Asp Ala Ile Lys Ser Ala Leu
177     210                 215                 220
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208 tggatatagg ggataaagaa accactgaaa gaatcgctct agcaagcggt cgtatcagca 300
210 tgaataaaga ggcttatgac gctattatca atcatggcgt caaaaagggt ccggtattac 360
212 aaactgctat tattgctggg attatggggg ctaaaaagac aagcgaactc attcccatgt 420
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231 Lys Cys Leu Leu Gly Ala Ser Val Val Ala Leu Leu Val Gly Cys Ser
232     15                  20                  25
234 ccg cat att att gaa acc aat gaa gtc gct ttg aaa ttg aat tac cat   927
235 Pro His Ile Ile Glu Thr Asn Glu Val Ala Leu Lys Leu Asn Tyr His
236  30                  35                  40                  45
238 cca gct agc gag aaa gtt caa gcg tta gat gaa aag att ttg ctt tta   975
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242 agg cca gct ttc caa tat agc gat aat atc gct aaa gag tat gaa aac  1023
243 Arg Pro Ala Phe Gln Tyr Ser Asp Asn Ile Ala Lys Glu Tyr Glu Asn
244              65                  70                  75
246 aaa ttc aag aat caa acc gcg ctc aag gtt gaa cag att ttg caa aat  1071
247 Lys Phe Lys Asn Gln Thr Ala Leu Lys Val Glu Gln Ile Leu Gln Asn
248          80                  85                  90
250 caa ggc tat aag gtt att agc gta gat agc agc gat aaa gac gat ttt  1119
251 Gln Gly Tyr Lys Val Ile Ser Val Asp Ser Ser Asp Lys Asp Asp Phe
252      95                  100                 105
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262 gaa ccc ggg tta tta ttc tcc acc ggt ttg gac aaa atg gaa ggg gtt  1263
263 Glu Pro Gly Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val
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272     175                 180                 185
274 caa gaa aaa ttc tta aaa acc acc cat tca agc cat agc ggg ggg tta  1407
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276 190                 195                 200                 205
278 gtt agc act atg gtt aag gga acg gat aat tct aat gac gcg atc aag  1455
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280                 210                 215                 220
282 aga gct ttg aat aag att ttt gca aat atc atg caa gaa ata gac aaa  1503
283 Arg Ala Leu Asn Lys Ile Phe Ala Asn Ile Met Gln Glu Ile Asp Lys
284              225                 230                 235
286 aaa ctc act caa aag aat tta gaa tct tat caa aaa gac gcc aaa gaa  1551
287 Lys Leu Thr Gln Lys Asn Leu Glu Ser Tyr Gln Lys Asp Ala Lys Glu
288          240                 245                 250
290 tta aaa ggc aaa aga aac cga taaaaacaaa taacgcataa gaaaagaacg    1602
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291 Leu Lys Gly Lys Arg Asn Arg
292      255             260
294 cttgaataaa ctgcttaaaa agggtttttt agcgttcttt ttgagcgtgt atttaagggc 1662
296 tgatgatc                                                            1670
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312             35              40                  45
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318  65                  70              75                  80
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321             85              90                  95
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324             100             105             110
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330     130             135             140
332 Leu Leu Phe Ser Thr Gly Leu Asp Lys Met Glu Gly Val Leu Ile Pro
333 145             150             155                 160
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336             165             170             175
338 Leu Asp Ser Phe Thr Met Asp Leu Ser Glu Leu Asp Ile Gln Glu Lys
339             180             185             190
341 Phe Leu Lys Thr Thr His Ser Ser His Ser Gly Gly Leu Val Ser Thr
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where are amino acids ?

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/308,435A**

DATE: 04/26/2001
TIME: 11:52:04

Input Set : **A:\Astra561.app**
Output Set: **N:\CRF3\04262001\I308435A.raw**

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